10/583061

HUI-054US

IAP20 Rec'd PCT/PTO 15 JUN 2006

SEQUENCE LISTING

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<110> Kerri MOWEN
      Laurie H. GLIMCHER
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745

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775

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                                                      30
cgt ttc ttc tat ccc gag ccg ggc gca cag gac ccg acc gat cgc cgc
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Arg Phe Phe Tyr Pro Glu Pro Gly Ala Gln Asp Pro Thr Asp Arg Arg
         35
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gca ggt agc agc ctg ggg acg ccc tac tct ggg ggc gcc ctg gtg cct
Ala Gly Ser Ser Leu Gly Thr Pro Tyr Ser Gly Gly Ala Leu Val Pro
                         55
gcc gcg ccg ggt cgc ttc ctt gga tcc ttc gcc tac ccg ccc cgg gct
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Ala Ala Pro Gly Arg Phe Leu Gly Ser Phe Ala Tyr Pro Pro Arg Ala
 65
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                                          75
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cag gtg gct ggc ttt ccc ggg cct ggc gag ttc ttc ccg ccg ccc gcg
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Gln Val Ala Gly Phe Pro Gly Pro Gly Glu Phe Phe Pro Pro Ala
                 85
                                      90
                                                          95
ggt gcg gag ggc tac ccg ccc gtg gat ggc tac cct gcc cct gac ccg
                                                                   336
Gly Ala Glu Gly Tyr Pro Pro Val Asp Gly Tyr Pro Ala Pro Asp Pro
            100
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cgc gcg ggg ctc tac cca ggg ccg cgc gag gac tac gca ttg ccc gcg
                                                                   384
Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro Ala
        115
                            120
                                                 125
ggg ttg gag gtg tct ggg aag ctg aga gtc gcg ctc agc aac cac ctg
                                                                   432
Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Ser Asn His Leu
    130
                                             140
                        135
ttg tgg tcc aag ttc aac cag cac cag aca gag atg atc atc act aag
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Leu 145	Trp	Ser	Lys	Phe	Asn 150	Gln	His	Gln	Thr	Glu 155	Met	Ile	Ile	Thr	Lys 160	
_	gga Gly		_	_				_					_		ctg Leu	528
	ccc Pro															576
	cac His															624
	gaa Glu 210							_			_			_		672
	aac Asn															720
	aag Lys													_	atg Met	768
	gtc Val															816
_	gtg Val							_	_	_	_	_				864
	gtc Val 290															912
	aac Asn															960
	gga Gly			Glu	Asn		Glu	Ser	Met	Tyr	Ala		Val		acg Thr	1008
	gtc Val														_	1056
	ttc Phe												_	_		1104
	ccc Pro 370															1152
	ctg Leu															1200

385	390	395	400								
	Thr Leu Leu Pro	tct gcc ccg ggg ccc act Ser Ala Pro Gly Pro Thr 410 415									
-		gcg cct gga gct ggt tgg Ala Pro Gly Ala Gly Trp 430									
		agc cca gct ggc tgg ttc Ser Pro Ala Gly Trp Phe 445									
		ggc ctg gga tcc tca gag Gly Leu Gly Ser Ser Glu 460									
		gag gtc acc tcc ctc cag Glu Val Thr Ser Leu Gln 475									
_	Gly Leu Gly Glu	gga gac act aag agg agg Gly Asp Thr Lys Arg Arg 490 495									
		agc tcc tct ccc gct ggg Ser Ser Ser Pro Ala Gly 510	_								
		ggc cag ttt tat aat tat Gly Gln Phe Tyr Asn Tyr 525									
ccc aac tga Pro Asn 530			1593								
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Pro Met Pro Ser Asp 20	Glu Gly Arg Gly 25	Pro Gly Ala Asp Gln Gln 30	His								
Arg Phe Phe Tyr Pro	Glu Pro Gly Ala (Gln Asp Pro Thr Asp Arg 45	Arg								
Ala Gly Ser Ser Leu 50	Gly Thr Pro Tyr 55	Ser Gly Gly Ala Leu Val 60	Pro								
Ala Ala Pro Gly Arg 65	Phe Leu Gly Ser 70	Phe Ala Tyr Pro Pro Arg 75	Ala 80								

Gln Val Ala Gly Phe Pro Gly Pro Gly Glu Phe Phe Pro Pro Pro Ala Gly Ala Glu Gly Tyr Pro Pro Val Asp Gly Tyr Pro Ala Pro Asp Pro Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro Ala Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Ser Asn His Leu Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr Lys Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly Leu Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val Asp Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly Lys Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp Ser Pro Asn Thr Gly Ala His Trp Met Arg Gln Glu Val Ser Phe Gly Lys Leu Lys Leu Thr Asn Asn Lys Gly Ala Ser Asn Asn Val Thr Gln Met Ile Val Leu Gln Ser Leu His Lys Tyr Gln Pro Arg Leu His Ile Val Glu Val Asn Asp Gly Glu Pro Glu Ala Ala Cys Ser Ala Ser Asn Thr - 285 His Val Phe Thr Phe Gln Glu Thr Gln Phe Ile Ala Val Thr Ala Tyr Gln Asn Ala Glu Ile Thr Gln Leu Lys Ile Asp Asn Asn Pro Phe Ala Lys Gly Phe Arg Glu Asn Phe Glu Ser Met Tyr Ala Ser Val Asp Thr Ser Val Pro Ser Pro Pro Gly Pro Asn Cys Gln Leu Leu Gly Gly Asp Pro Phe Ser Pro Leu Leu Ser Asn Gln Tyr Pro Val Pro Ser Arg Phe Tyr Pro Asp Leu Pro Gly Gln Pro Lys Asp Met Ile Ser Gln Pro Tyr Trp Leu Gly Thr Pro Arg Glu His Ser Tyr Glu Ala Glu Phe Arg Ala

Val Ser Met Lys Pro Thr Leu Leu Pro Ser Ala Pro Gly Pro Thr Val 405 410 415

Pro Tyr Tyr Arg Gly Gln Asp Val Leu Ala Pro Gly Ala Gly Trp Pro 420 425 430

Val Ala Pro Gln Tyr Pro Pro Lys Met Ser Pro Ala Gly Trp Phe Arg 435 440 445

Pro Met Arg Thr Leu Pro Met Asp Pro Gly Leu Gly Ser Ser Glu Glu 450 455 460

Gln Gly Ser Ser Pro Ser Leu Trp Pro Glu Val Thr Ser Leu Gln Pro 465 470 475 480

Glu Pro Ser Asp Ser Gly Leu Gly Glu Gly Asp Thr Lys Arg Arg Arg 495

Ile Ser Pro Tyr Pro Ser Ser Gly Asp Ser Ser Ser Pro Ala Gly Ala 500 505 510

Pro Ser Pro Phe Asp Lys Glu Thr Glu Gly Gln Phe Tyr Asn Tyr Phe 515 520 525

Pro Asn 530